HS_sponge_AvN_names

| All v records per page | Search all columns: | | | |
|--|--|--------|-------------|------------------|
| | | From | From | From |
| | | to | to | to |
| | | | | |
| | Image | logFC | p- Value | Adjusted p-Value |
| ogroup8371_c0_g1_NA | 6 | -28.20 | | 1.96e-03 |
| ogroup30126_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8371_c0_g1_NA.pdf) | -27.60 | 3.38e-06 | 2.80e-03 |
| ogroup4139_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup30126_c0_g1_NA.pdf) | -27.50 | 3.77e-06 | 2.94e-03 |
| ogreupg | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4139_c0_g1_NA.pdf) | 27.00 | 070 00 | 2.0 10 00 |
| ogroup15081_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15081_c0_g1_NA.pdf) | -26.30 | 9.95e-06 | 5.94e-03 |
| ogroup7412_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7412_c0_g2_NA.pdf) | -25.90 | 1.38e-05 | 7.23e-03 |
| ogroup7412_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7412_c0_g1_NA.pdf) | -25.20 | 2.33e-05 | 1.04e-02 |
| ogroup1399_c0_g1_FTH1_oxidoreductase_activityo | b (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup1399_c0_g1_FTH1 | -25.10 | 2.47e-05 | 1.05e-02 |
| ogroup30031_c0_g3_PCK1_phosphoenolpyruvate_carbo | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup30031_c0_g3_PCK | -25.10 | 2.41e-05 | 1.04e-02 |
| ogroup5983_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5983_c0_g2_NA.pdf) | -25.10 | 2.39e-05 | 1.04e-02 |
| ogroup5843_c0_g1_EZR_protein_localization_to_cel | | -25.00 | 2.72e-05 | 1.09e-02 |
| ogroup23156_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup23156_c0_g1_NA.pdf) | -24.70 | 3.29e-05 | 1.26e-02 |
| ogroup19142_c0_g1_TPT1_calcium_ion_binding | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19142_c0_g1_TPT1 | -24.60 | 3.64e-05 | 1.35e-02 |
| ogroup5977_c0_g1_RPL6_60s_ribosomal_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5977_c0_g1_RPL6 | -24.60 | 3.44e-05 | 1.30e-02 |
| ogroup4139_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4139_c0_g2_NA.pdf) | -24.50 | 3.70e-05 | 1.36e-02 |
| ogroup10714_c0_g1_RPL21_ribosomal_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10714_c0_g1_RPL | -24.40 | 4.24e-05 | 1.47e-02 |
| ogroup7295_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7295_c0_g1_NA.pdf) | -24.40 | 4.21e-05 | 1.47e-02 |
| ogroup22202_c0_g3_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup22202_c0_g3_NA.pdf) | -24.30 | 1.61e-09 | 4.86e-06 |
| ogroup2762_c0_g5_HSP90AA1_Molecular_chaperone_th | • • • • • • • • • • • • • • • • • • • | -24.30 | 4.42e-05 | 1.49e-02 |
| ogroup3650_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3650_c0_g1_NA.pdf) | -24.30 | 4.44e-05 | 1.49e-02 |
| ogroup6430_c0_g1_NA | • ○ (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6430_c0_g1_NA.pdf) | -24.30 | 4.26e-05 | 1.47e-02 |
| ogroup6883_c0_g1_RPL18_structural_constituent_of | | -24.20 | 4.71e-05 | 1.54e-02 |
| ogroup6450_c0_g1C-type_lectinCTLor_carbohy | • (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6450_c0_g1C-typ | -24.10 | 4.94e-05 | 1.56e-02 |
| ogroup6820_c0_g1_NA | • • • • • • • • • • • • • • • • • • • | -24.10 | 4.97e-05 | 1.56e-02 |
| ogroup9712_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9712_c0_g2_NA.pdf) | -24.00 | 5.38e-05 | 1.67e-02 |

| | | From | From | From |
|--|--|--------|-------------|------------------|
| | | to | to | to |
| ID | Image | logFC | p- Value | Adjusted p-Value |
| isogroup10430_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10430_c0_g1_NA.pdf) | -23.90 | 5.68e-05 | 1.70e-02 |
| isogroup7050_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7050_c0_g1_NA.pdf) | -23.90 | 5.68e-05 | 1.70e-02 |
| isogroup13506_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13506_c0_g1_NA.pdf) | -23.80 | 6.53e-05 | 1.79e-02 |
| isogroup163042_c0_g1_GNB2L1_positive_regulation_of | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup163042_c0_g1_GN | -23.80 | 6.22e-05 | 1.78e-02 |
| isogroup6982_c0_g2Chitin-binding_domain_type_2 | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6982_c0_g2Chiti | -23.80 | 6.19e-05 | 1.78e-02 |
| isogroup7099_c0_g1translocon-associated_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7099_c0_g1transl | -23.80 | 6.14e-05 | 1.78e-02 |
| isogroup85143_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup85143_c0_g2_NA.pdf) | -23.80 | 6.27e-05 | 1.78e-02 |
| isogroup88347_c0_g1_RPS2_translation | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup88347_c0_g1_RPS | -23.80 | 6.37e-05 | 1.79e-02 |
| isogroup7742_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7742_c0_g2_NA.pdf) | -23.70 | 6.67e-05 | 1.81e-02 |
| isogroup8501_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8501_c0_g1_NA.pdf) | -23.70 | 6.92e-05 | 1.82e-02 |
| isogroup25235_c0_g2_NA | 6 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup25235_c0_g2_NA.pdf) | -23.60 | 7.50e-05 | 1.87e-02 |
| isogroup26694_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup26694_c0_g1_NA.pdf) | -23.60 | 7.40e-05 | 1.87e-02 |
| isogroup51929_c0_g3_RpL3_ribosomal_large_subunit_a | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup51929_c0_g3_RpL3 | -23.60 | 7.29e-05 | 1.86e-02 |
| isogroup1399_c0_g2_RpL11_Belongs_to_the_universal_ | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup1399_c0_g2_RpL11 | -23.50 | 7.81e-05 | 1.91e-02 |
| isogroup19842_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19842_c0_g1_NA.pdf) | -23.50 | 7.97e-05 | 1.93e-02 |
| isogroup15032_c0_g1_HSP90B1_heat_shock_protein_90k | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15032_c0_g1_HSP | -23.40 | 8.30e-05 | 1.98e-02 |
| isogroup43972_c0_g4_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup43972_c0_g4_NA.pdf) | -23.40 | 8.44e-05 | 1.98e-02 |
| isogroup48008_c1_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup48008_c1_g1_NA.pdf) | -23.40 | 8.65e-05 | 1.99e-02 |
| isogroup57530_c0_g1_rps15a_structural_constituent_ | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup57530_c0_g1_rps1 | -23.40 | 8.34e-05 | 1.98e-02 |
| isogroup13958_c0_g2Fibrinogen-related_domainsF | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13958_c0_g2_Fibr | -23.30 | 8.87e-05 | 2.02e-02 |
| isogroup21068_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup21068_c0_g1_NA.pdf) | -23.30 | 2.28e-09 | 5.51e-06 |
| isogroup112313_c0_g1_FOLR1_Folate_receptor | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup112313_c0_g1_FOL | -23.20 | 9.47e-05 | 2.09e-02 |
| isogroup11804_c0_g1Calmodulin | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup11804_c0_g1Cal | -23.20 | 9.85e-05 | 2.09e-02 |
| isogroup126047_c0_g1actin_binding | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup126047_c0_g1act | -23.20 | 9.53e-05 | 2.09e-02 |
| isogroup15465_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15465_c0_g1_NA.pdf) | -23.20 | 9.57e-05 | 2.09e-02 |
| isogroup21023_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup21023_c0_g1_NA.pdf) | -23.20 | 9.88e-05 | 2.09e-02 |

| | | From | From | From |
|---|--|------------------|----------|----------------------|
| | | to | to | to |
| | | | p- | Adjusted |
| ID | Image | logFC | Value | p-Value |
| isogroup6350_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6350_c0_g1_NA.pdf) | -23.20 | 9.57e-05 | 2.09e-02 |
| isogroup6353_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6353_c0_g1_NA.pdf) | -23.20 | 9.94e-05 | 2.09e-02 |
| isogroup9062_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9062_c0_g1_NA.pdf) | -23.20 | 9.39e-05 | 2.09e-02 |
| isogroup9804_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9804_c0_g2_NA.pdf) | -23.20 | 9.92e-05 | 2.09e-02 |
| isogroup12003_c0_g4_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12003_c0_g4_NA.pdf) | -23.00 | 1.08e-04 | 2.10e-02 |
| isogroup14392_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14392_c0_g1_NA.pdf) | -23.00 | 1.08e-04 | 2.10e-02 |
| isogroup65255_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup65255_c0_g2_NA.pdf) | -23.00 | 1.13e-04 | 2.10e-02 |
| isogroup7826_c0_g1_RPLP0_ribosome_biogenesis | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7826_c0_g1_RPLP | -23.00 | 1.11e-04 | 2.10e-02 |
| isogroup8328_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8328_c0_g1_NA.pdf) | -23.00 | 1.13e-04 | 2.10e-02 |
| isogroup10269_c0_g1_dll3_Delta-like_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10269_c0_g1_dll3 | -22.90 | 1.20e-04 | 2.10e-02 |
| isogroup104477_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup104477_c0_g1_NA | -22.90 | 1.17e-04 | 2.10e-02 |
| isogroup112646_c0_g1_SEC61G_Protein_transport_prot | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup112646_c0_g1_SE | -22.90 | 1.18e-04 | 2.10e-02 |
| isogroup14208_c0_g1_SSR4_Translocon-associated_pro | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14208_c0_g1_SSR | -22.90 | | 2.10e-02 |
| isogroup15723_c1_g3_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15723_c1_g3_NA.pdf) | -22.90 | 1.20e-04 | 2.10e-02 |
| isogroup16167_c0_g1neurogenic_locus_notch_homolo | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16167_c0_g1_neu | -22.90 | | 2.10e-02 |
| isogroup17735_c0_g1Fibrinogen-related_domainsF | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup17735_c0_g1Fibr | | | 2.10e-02 |
| isogroup26354_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup26354_c0_g1_NA.pdf) | | | 2.10e-02 |
| isogroup5983_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5983_c0_g1_NA.pdf) | -22.90 -22.90 | | 2.10e-02 2.10e-02 |
| isogroup6289_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6289_c0_g1_NA.pdf) | | 1.18e-04 | |
| isogroup7222_c0_g1_NA isogroup7243_c0_g1_RPS25_ribosomal_protein_S25 | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7222_c0_g1_NA.pdf) | | 1.18e-04 | |
| isogroup8063_c0_g2_eef1g_translation_elongation_fa | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7243_c0_g1_RPS2 | -22.90 | | 2.10e-02 |
| isogroup8375 c0 g2 actin filament capping | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8063_c0_g2_eef1g | -22.90 | | 2.10e-02 |
| isogroup9002_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8375_c0_g2_actin | | | 2.10e-02 |
| isogroup9826_c0_g1_MDH1_Malate_dehydrogenase | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9002_c0_g2_NA.pdf) | -22.90 | | 2.10e-02 |
| isogroup10936_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9826_c0_g1_MDH1 | -22.80 | | 3.41e-06 |
| | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10936_c0_g2_NA.pdf) | | | |

From

| to | | to | From |
|--|---------|-------------|------------------|
| · | 0 | | |
| ID Image | | p- Value | Adjusted p-Value |
| isogroup11243_c0_g1_NA [figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup11243_c0_g1_NA.pdf] -2 | 22.80 1 | 1.26e-04 | 2.14e-02 |
| isogroup18712_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup18712_c0_g1_NA.pdf) | 22.80 1 | 1.29e-04 | 2.18e-02 |
| isogroup19447_c0_g1_HSP90B1_heat_shock_protein_90k (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19447_c0_g1_HSP2 | 22.80 1 | 1.30e-04 | 2.18e-02 |
| isogroup10381_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10381_c0_g1_NA.pdf) | 22.70 1 | 1.38e-04 | 2.18e-02 |
| isogroup11495_c0_g2_NA [figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup11495_c0_g2_NA.pdf] -2 | 22.70 1 | 1.36e-04 | 2.18e-02 |
| isogroup13585_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13585_c0_g1_NA.pdf) | 22.70 1 | 1.38e-04 | 2.18e-02 |
| isogroup2475_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup2475_c0_g1_NA.pdf) | 22.70 1 | 1.37e-04 | 2.18e-02 |
| isogroup32998_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup32998_c0_g1_NA.pdf) -2 | 22.70 1 | 1.34e-04 | 2.18e-02 |
| isogroup5853_c0_g3_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5853_c0_g3_NA.pdf) | 22.70 1 | 1.38e-04 | 2.18e-02 |
| isogroup9002_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9002_c0_g1_NA.pdf) | 22.70 1 | 1.36e-04 | 2.18e-02 |
| isogroup9467_c0_g2_GTPBP4_Involved_in_the_biogenes (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9467_c0_g2_GTPB2 | 22.70 1 | 1.39e-04 | 2.18e-02 |
| isogroup10920_c0_g3_RPL27A_ribosomal_protein (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10920_c0_g3_RPL2 | 22.60 1 | 1.44e-04 | 2.18e-02 |
| isogroup14452_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14452_c0_g1_NA.pdf) -2 | 22.60 1 | 1.45e-04 | 2.18e-02 |
| isogroup19367_c0_g1_NA | 22.60 1 | 1.43e-04 | 2.18e-02 |
| isogroup20229_c0_g2_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup20229_c0_g2_NA.pdf) -2 | 22.60 1 | 1.48e-04 | 2.18e-02 |
| isogroup23677_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup23677_c0_g1_NA.pdf) | 22.60 1 | 1.43e-04 | 2.18e-02 |
| isogroup4098_c0_g4_meprin_and_TRAF_homology | 22.60 1 | 1.96e-05 | 9.28e-03 |
| isogroup78210_c0_g1_NA | 22.60 1 | 1.48e-04 | 2.18e-02 |
| isogroup8750_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8750_c0_g1_NA.pdf) | 22.60 1 | 1.41e-04 | 2.18e-02 |
| isogroup10341_c0_g2_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10341_c0_g2_NA.pdf) -2 | 22.50 1 | 1.50e-04 | 2.18e-02 |
| isogroup12499_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12499_c0_g1_NA.pdf) -2 | 22.50 1 | 1.55e-04 | 2.19e-02 |
| isogroup13868_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13868_c0_g1_NA.pdf) -2 | 22.50 1 | 1.54e-04 | 2.19e-02 |
| isogroup15723_c1_g4_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15723_c1_g4_NA.pdf) | 22.50 1 | 1.51e-04 | 2.18e-02 |
| | 22.50 1 | 1.62e-08 | 2.78e-05 |
| isogroup6832_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6832_c0_g1_NA.pdf) | 22.50 1 | 1.51e-04 | 2.18e-02 |
| isogroup81675_c0_g1_NA | 22.50 1 | 1.50e-04 | 2.18e-02 |

| | | From | From to | From to |
|--|--|--------|----------|------------|
| | | to | | |
| | | | p- | Adjusted |
| ID | Image | logFC | Value | p-Value |
| isogroup83577_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup83577_c0_g1_NA.pdf) | -22.50 | 1.52e-04 | 2.19e-02 |
| isogroup8807_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8807_c0_g1_NA.pdf) | -22.50 | 1.54e-04 | 2.19e-02 |
| isogroup15508_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15508_c0_g1_NA.pdf) | -22.40 | 1.63e-04 | 2.22e-02 |
| isogroup15617_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15617_c0_g1_NA.pdf) | -22.40 | 1.65e-04 | 2.22e-02 |
| isogroup25392_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup25392_c0_g1_NA.pdf) | -22.40 | 1.70e-04 | 2.24e-02 |
| isogroup4596_c0_g2_RPLP1_protein_kinase_activator_ | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4596_c0_g2_RPLP | -22.40 | 1.66e-04 | 2.22e-02 |
| isogroup6979_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6979_c0_g1_NA.pdf) | -22.40 | 1.61e-04 | 2.22e-02 |
| isogroup7673_c0_g3_eef2_Elongation_factor | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7673_c0_g3_eef2 | -22.40 | 1.67e-04 | 2.22e-02 |
| isogroup8747_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8747_c0_g1_NA.pdf) | -22.40 | 1.64e-04 | 2.22e-02 |
| isogroup9467_c0_g3_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9467_c0_g3_NA.pdf) | -22.30 | 1.89e-09 | 5.08e-06 |
| isogroup13626_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13626_c0_g1_NA.pdf) | -22.20 | 3.57e-08 | 5.08e-05 |
| isogroup112876_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup112876_c0_g1_NA | -22.10 | 2.02e-04 | 2.53e-02 |
| isogroup12025_c0_g1_PDAP1_28_kDa_heatand_acid-st | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12025_c0_g1_PDA | -22.10 | 2.01e-04 | 2.53e-02 |
| isogroup51601_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup51601_c0_g2_NA.pdf) | -22.10 | 1.97e-04 | 2.49e-02 |
| isogroup7501_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7501_c0_g2_NA.pdf) | -22.10 | 3.18e-08 | 4.80e-05 |
| isogroup11246_c0_g1_STT3A_co-translational_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup11246_c0_g1_STT3 | -22.00 | 2.13e-04 | 2.66e-02 |
| isogroup34916_c0_g1dolichyl-diphosphooligosaccha | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup34916_c0_g1_doli | -22.00 | 2.21e-04 | 2.74e-02 |
| isogroup32001_c0_g4_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup32001_c0_g4_NA.pdf) | -21.90 | 8.91e-09 | 1.80e-05 |
| isogroup4376_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4376_c0_g2_NA.pdf) | -21.90 | 2.30e-04 | 2.79e-02 |
| isogroup8251_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8251_c0_g1_NA.pdf) | -21.90 | 2.28e-04 | 2.79e-02 |
| isogroup12563_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12563_c0_g1_NA.pdf) | -21.80 | 1.35e-08 | 2.51e-05 |
| isogroup4376_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4376_c0_g1_NA.pdf) | -21.80 | 2.40e-04 | 2.87e-02 |
| isogroup56570_c0_g1_NA | | -21.80 | 1.81e-11 | 1.09e-07 |
| isogroup3286_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3286_c0_g2_NA.pdf) | -21.70 | 2.72e-04 | 3.24e-02 |
| isogroup9393_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9393_c0_g1_NA.pdf) | -21.50 | 3.01e-04 | 3.54e-02 |
| isogroup5162_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5162_c0_g1_NA.pdf) | -21.40 | 3.20e-04 | 3.74e-02 |
| | | | | |

From

| | | From | From | From |
|--|---|--------|-------------|------------------|
| | | to | to | to |
| | | | | |
| ID | Image | logFC | p- Value | Adjusted p-Value |
| isogroup9831_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9831_c0_g1_NA.pdf) | -21.20 | | 4.04e-02 |
| isogroup9522_c0_g1activation_of_NF-kappaB-induci | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9522_c0_g1activ | -21.00 | 4.27e-04 | 4.61e-02 |
| isogroup9792_c0_g2_ATP6V0C_Proton-conducting_pore_ | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9792_c0_g2_ATP6 | -20.50 | 5.77e-04 | 5.82e-02 |
| isogroup17629_c0_g4_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup17629_c0_g4_NA.pdf) | -20.40 | 3.90e-04 | 4.33e-02 |
| isogroup3478_c0_g1_FAM83H_keratin_filament_binding | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3478_c0_g1_FAM8 | -20.00 | 7.59e-04 | 7.12e-02 |
| isogroup9440_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9440_c0_g1_NA.pdf) | -20.00 | 7.88e-04 | 7.30e-02 |
| isogroup11653_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup11653_c0_g1_NA.pdf) | -19.80 | 8.54e-04 | 7.74e-02 |
| isogroup84941_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup84941_c0_g2_NA.pdf) | -19.20 | 3.48e-06 | 2.80e-03 |
| isogroup10077_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10077_c0_g2_NA.pdf) | -17.20 | 3.22e-04 | 3.74e-02 |
| isogroup10745_c0_g1_EF1a-F2_regulation_of_response | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10745_c0_g1_EF1a | -15.40 | 4.25e-06 | 3.03e-03 |
| isogroup17582_c0_g2_FTH1_oxidoreductase_activity | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup17582_c0_g2_FTH | -14.90 | 1.22e-10 | 5.88e-07 |
| isogroup14921_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14921_c0_g1_NA.pdf) | -14.50 | 8.62e-05 | 1.99e-02 |
| isogroup13089_c0_g1_RPLP2_Belongs_to_the_eukaryoti | b (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13089_c0_g1_RPL b - ◆ - ◆ - ◆ - ◆ - ◆ - | -14.20 | 4.82e-05 | 1.56e-02 |
| isogroup10957_c0_g1_NA | b (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10957_c0_g1_NA.pdf) c (\$\sqrt{2}\ | -14.10 | 4.09e-06 | 3.00e-03 |
| isogroup15873_c0_g2_NA | 6 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15873_c0_g2_NA.pdf) | -13.50 | 1.86e-04 | 2.42e-02 |
| isogroup949_c1_g1_NA | | -13.50 | 9.56e-10 | 3.41e-06 |
| isogroup3910_c1_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3910_c1_g1_NA.pdf) | -13.40 | 7.71e-05 | 1.90e-02 |
| isogroup56436_c1_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup56436_c1_g1_NA.pdf) | -13.40 | 1.46e-04 | 2.18e-02 |
| isogroup7246_c0_g6protein_modification_by_small_ | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7246_c0_g6_prote | | 1.97e-04 | |
| isogroup9722_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9722_c0_g1_NA.pdf) | | 3.04e-05 | |
| isogroup12581_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12581_c0_g1_NA.pdf) | | 4.53e-05 | |
| isogroup12789_c1_g1BPTI_Kunitz_family_of_serine_ | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12789_c1_g1BP | | 3.14e-09 | |
| | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup114811_c0_g1_RPS | | 6.74e-05 | |
| isogroup7119_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7119_c0_g1_NA.pdf) | | 2.87e-04 | |
| isogroup7785_c0_g1_RPL7_ribosomal_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7785_c0_g1_RPL7 | | | |
| isogroup95748_c0_g1_RpS27A_ubiquitin | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup95748_c0_g1_RpS | -12.80 | 5.76e-04 | 5.82e-02 |
| | | | | |

| ID Image Isogroup33375_c1_g1_RPL35A_structural_constituent_ (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup33375_c1_g1_RPL12.4 | 40 1.69e-04 40 8.61e-08 10 2.92e-04 00 7.39e-04 30 1.04e-03 30 1.22e-05 | Adjusted p-Value 2.23e-02 1.16e-04 3.45e-02 7.04e-02 9.17e-02 6.68e-03 |
|--|--|--|
| isogroup33375_c1_g1_RPL35A_structural_constituent | FC Value 40 1.69e-04 40 8.61e-08 10 2.92e-04 00 7.39e-04 30 1.04e-03 30 1.22e-05 | p-Value 2.23e-02 1.16e-04 3.45e-02 7.04e-02 9.17e-02 |
| isogroup3547_c0_g3_NA isogroup4609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup3113_c1_g3_NA isogroup3113_c1_g3_NA isogroup3547_c0_g1_NAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4609_c0_g1_RPS1 isogroup7246_c0_g5_NA isogroup3113_c1_g3_NA isogroup3113_c1_g3_NA isogroup3547_c0_g1_SFRNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7246_c0_g5_NA.pdf) -12.0 -12 | 40 8.61e-08 10 2.92e-04 00 7.39e-04 30 1.04e-03 30 1.22e-05 | 1.16e-04 3.45e-02 7.04e-02 9.17e-02 |
| isogroup3547_c0_g3_NA isogroup4609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup3113_c1_g3_NA isogroup3547_c0_g3_NA isogroup3113_c1_g3_NA isogroup34609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup7246_c0_g5_NA isogroup3113_c1_g3_NA isogroup34609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup7246_c0_g5_NA isogroup7246_c0_g5_NA isogroup34609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup7246_c0_g5_NA isogroup7246_c0_g5_NA isogroup34609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup7246_c0_g5_NA isogroup7246_c0_g5_NA isogroup34609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup72 | 10 2.92e-04 00 7.39e-04 30 1.04e-03 1.22e-05 | 3.45e-02 7.04e-02 9.17e-02 |
| isogroup4609_c0_g1_RPS16_Belongs_to_the_universal_ isogroup7246_c0_g5_NA isogroup3113_c1_g3_NA isogroup9547_c0_g1_SS_rRNA binding isogroup9547_c0_g1_SS_rRNA binding | 7.39e-04 30 1.04e-03 30 1.22e-05 | 7.04e-02 9.17e-02 |
| isogroup3113_c1_g3_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3113_c1_g3_NA.pdf) isogroup9547_c0_g1_5S_rRNA_binding | 30 1.04e-03 30 1.22e-05 | 9.17e-02 |
| isogroup9547 c0 g1 5S rRNA binding | 30 1.22e-05 | |
| isogroup9547_c0_g15S_rRNA_binding | | 6.68e-03 |
| · · · · · · · · · · · · · · · · · · · | 70 7.55e-07 | |
| isogroup19836_c1_g1_NA | | 8.70e-04 |
| isogroup13429_c0_g1_zinc_ion_binding | 9.60e-07 | 1.01e-03 |
| isogroup15723_c1_g2_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup15723_c1_g2_NA.pdf) -11.6 | 30 8.04e-06 | 5.28e-03 |
| isogroup19650_c0_g1_serine-type_endopeptidase_act (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19650_c0_g1_seri11.6 | 30 1.23e-04 | 2.12e-02 |
| isogroup8001_c0_g2_ENO3_Enolase_3betamuscle_ (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8001_c0_g2_ENO3 | 30 5.21e-04 | 5.34e-02 |
| isogroup9860_c1_g1_NA | i0 2.57e-05 | 1.07e-02 |
| isogroup22322_c0_g3_RPS7_structural_constituent_of (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup22322_c0_g3_RPS (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup22322_c0_g3_RPS | 10 2.26e-04 | 2.78e-02 |
| isogroup24667_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup24667_c0_g1_NA.pdf) | 20 1.12e-04 | 2.10e-02 |
| isogroup292_c0_g2_NA | 10 4.08e-13 | 3.29e-09 |
| isogroup43264_c0_g1_NA | 10 4.72e-04 | 5.02e-02 |
| isogroup43972_c0_g3_NA | 1.35e-04 | 2.18e-02 |
| isogroup56345_c0_g2_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup56345_c0_g2_NA.pdf) | 0 8.08e-04 | 7.44e-02 |
| isogroup8372_c1_g1_NA | 10 1.24e-05 | 6.68e-03 |
| isogroup7246_c0_g3_NA | 0 9.07e-04 | 8.16e-02 |
| isogroup19380_c0_g2_NA | 90 1.81e-05 | 8.94e-03 |
| isogroup4829_c0_g2_RPS2_ribosomal_protein_S2 | 80 7.86e-06 | 5.28e-03 |
| isogroup586_c7_g1_eef2_translation_elongation_fact | 80 4.00e-04 | 4.41e-02 |
| isogroup55102_c0_g1_rps9_positive_regulation_of_tr | 70 6.97e-04 | 6.77e-02 |
| isogroup10352_c0_g2_GTPBP2_GTPase_activity | 3.99e-06 | 3.00e-03 |
| isogroup16595_c1_g1_AIFM1_cellular_response_to_ald (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16595_c1_g1_AIFM (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16595_c1_g1_AIFM | 60 3.26e-04 | 3.76e-02 |

| | | FIOIII | From | From |
|--|--|--------|----------|----------|
| | | to | το | to |
| | | | p- | Adjusted |
| ID | Image | logFC | Value | p-Value |
| isogroup3057_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3057_c0_g1_NA.pdf) | -10.50 | 1.12e-03 | 9.69e-02 |
| isogroup249_c0_g1_RPL24_ribosomal_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup249_c0_g1_RPL24 | -10.40 | 7.66e-16 | 1.85e-11 |
| isogroup4118_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4118_c0_g1_NA.pdf) | -10.40 | 1.03e-03 | 9.09e-02 |
| isogroup7039_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7039_c0_g1_NA.pdf) | -10.40 | 7.48e-04 | 7.07e-02 |
| isogroup7414_c0_g2_NA | | -10.20 | 9.81e-06 | 5.94e-03 |
| isogroup11619_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup11619_c0_g2_NA.pdf) | -10.10 | 7.11e-05 | 1.84e-02 |
| isogroup3778_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3778_c0_g1_NA.pdf) | -9.89 | 1.10e-04 | 2.10e-02 |
| isogroup13746_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13746_c0_g1_NA.pdf) | -9.72 | 3.72e-04 | 4.15e-02 |
| isogroup8267_c0_g1_RPS5_ribosomal_small_subunit_as | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8267_c0_g1_RPS5 | -9.71 | 4.80e-04 | 5.08e-02 |
| isogroup34241_c0_g2_NA | | -9.68 | 4.02e-05 | 1.45e-02 |
| isogroup19424_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19424_c0_g1_NA.pdf) | -9.49 | 3.24e-04 | 3.75e-02 |
| isogroup16449_c0_g1_RPS3_positive_regulation_of_DN | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16449_c0_g1_RPS | -9.46 | 1.61e-04 | 2.22e-02 |
| isogroup3483_c0_g2_pdzrn4_PDZ_domain_containing_ri | | -9.45 | 5.48e-05 | 1.68e-02 |
| isogroup5557_c0_g1_rpl26l1_regulation_of_translati | | -9.42 | 2.31e-04 | 2.79e-02 |
| isogroup13716_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13716_c0_g1_NA.pdf) | -9.38 | 1.58e-05 | 8.13e-03 |
| isogroup6173_c0_g1_RPL13_structural_constituent_of | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup6173_c0_g1_RPL1 | -9.34 | 1.09e-04 | 2.10e-02 |
| isogroup39689_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup39689_c0_g1_NA.pdf) | -9.25 | 6.47e-05 | 1.79e-02 |
| isogroup9960_c1_g2Molecular_calcium-binding_chap | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9960_c1_g2_Mole | -9.12 | 9.19e-04 | 8.23e-02 |
| isogroup58989_c0_g1_RPL12_ribosomal_large_subunit_ | | -9.08 | 2.38e-04 | 2.86e-02 |
| isogroup9285_c2_g1NAD_P_H_oxidase_activity | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9285_c2_g1NAD | -9.05 | 1.81e-04 | |
| isogroup14560_c0_g1NAD_NADP_octopine_nopaline_de | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14560_c0_g1_NA | -9.04 | 5.17e-04 | |
| isogroup245_c0_g2G_U_mismatch-specific_uracil-DN | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup245_c0_g2G_U | -8.97 | 6.01e-04 | |
| isogroup45872_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup45872_c0_g1_NA.pdf) | -8.90 | 4.56e-04 | |
| isogroup20497_c0_g5_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup20497_c0_g5_NA.pdf) | | 1.02e-03 | |
| isogroup9954_c0_g1_RPL23_large_ribosomal_subunit_r | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9954_c0_g1_RPL2 | -8.78 | 6.90e-05 | |
| isogroup16997_c0_g3_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16997_c0_g3_NA.pdf) | -8.70 | 3.51e-04 | 4.01e-02 |
| | | | | |

| | | FIOIII | From | From |
|--|--|--------|-------------|------------------|
| | | to | | |
| ID | Image | logFC | p- Value | Adjusted p-Value |
| isogroup34544_c0_g3_RAB14_Golgi_to_endosome_transp | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup34544_c0_g3_RAB | -8.60 | 8.09e-04 | |
| isogroup10143_c0_g1_NA | o o o o o o o o o o o o o o o o o o o | -8.55 | 1.14e-03 | 9.76e-02 |
| isogroup24212_c3_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup24212_c3_g1_NA.pdf) | -8.46 | 6.87e-04 | 6.70e-02 |
| isogroup98537_c2_g2_TMSB4X_Thymosin_beta-4_family | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup98537_c2_g2_TMS | -8.44 | 1.15e-04 | 2.10e-02 |
| isogroup45542_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup45542_c0_g2_NA.pdf) | -8.30 | 6.48e-04 | 6.43e-02 |
| isogroup4841_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4841_c0_g2_NA.pdf) | -8.08 | 2.01e-05 | 9.35e-03 |
| isogroup13113_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13113_c0_g1_NA.pdf) | -8.06 | 4.45e-07 | 5.38e-04 |
| isogroup5273_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5273_c0_g1_NA.pdf) | -8.04 | 7.16e-04 | 6.93e-02 |
| isogroup24207_c1_g1cytoskeleton_organization | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup24207_c1_g1_cyto | -7.89 | 8.57e-04 | 7.74e-02 |
| isogroup12942_c2_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12942_c2_g1_NA.pdf) | -7.84 | 8.07e-06 | 5.28e-03 |
| isogroup4016_c0_g1_RPS12_structural_constituent_of | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup4016_c0_g1_RPS1 | -7.84 | 1.44e-04 | 2.18e-02 |
| isogroup2448_c0_g5_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup2448_c0_g5_NA.pdf) | -7.78 | 1.13e-03 | 9.69e-02 |
| isogroup13975_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13975_c0_g1_NA.pdf) | -7.72 | 4.41e-04 | 4.74e-02 |
| isogroup18880_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup18880_c0_g1_NA.pdf) | -7.67 | 8.57e-04 | 7.74e-02 |
| isogroup23870_c0_g4_NA | | -7.65 | 1.19e-07 | 1.52e-04 |
| isogroup43858_c0_g1_DBNL_podosome_assembly | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup43858_c0_g1_DBN | -7.62 | 8.49e-04 | 7.74e-02 |
| isogroup14912_c0_g3Collagen_triple_helix | • (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14912_c0_g3_Coll | -7.52 | 1.06e-03 | 9.19e-02 |
| isogroup16599_c0_g1_RPL17_structural_constituent_o | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16599_c0_g1_RPL | -7.52 | 1.65e-04 | 2.22e-02 |
| isogroup41535_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup41535_c0_g2_NA.pdf) | -7.51 | 1.73e-08 | 2.78e-05 |
| isogroup949_c0_g2_SMOC1_SPARC_related_modular_calc | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup949_c0_g2_SMOC1 | -7.45 | 3.57e-13 | 3.29e-09 |
| isogroup38426_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup38426_c0_g1_NA.pdf) | -7.43 | 1.16e-03 | |
| isogroup134495_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup134495_c0_g1_NA | -7.39 | 4.99e-04 | |
| isogroup14325_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14325_c0_g2_NA.pdf) | -6.82 | 1.17e-05 | |
| isogroup26070_c0_g2_ATP5A1_proton-transporting_ATP | | -6.39 | 2.78e-06 | |
| isogroup19804_c0_g2_PABPC1_Poly_A_binding_protein_ | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19804_c0_g2_PAB | -6.10 | 5.06e-04 | |
| isogroup19724_c1_g1_RPL5_ribosomal_protein | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19724_c1_g1_RPL | -6.04 | 1.01e-05 | 5.94e-03 |
| | | | | |

| | | From | From | From |
|--|--|-------|-------------|------------------|
| | | to | to | to |
| | | | | |
| ID | Image | logEC | p- Value | Adjusted p-Value |
| isogroup2422_c0_g3_Zinc_fingerC4_typetwo_doma | Image (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup2422_c0_g3_Zinc | -5.67 | | 1.84e-02 |
| isogroup31310_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup31310_c0_g2_NA.pdf) | -5.49 | 2.72e-06 | 2.40e-03 |
| isogroup13123_c0_g1_EEF1D_Belongs_to_the_EF-1-beta | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13123_c0_g1_EEF | -5.45 | 1.05e-03 | 9.18e-02 |
| isogroup9401_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9401_c0_g1_NA.pdf) | -5.41 | 1.05e-03 | 9.18e-02 |
| isogroup18121_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup18121_c0_g1_NA.pdf) | -5.15 | 3.56e-04 | 4.03e-02 |
| isogroup11619_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup11619_c0_g1_NA.pdf) | -5.13 | 1.89e-04 | 2.44e-02 |
| isogroup14428_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14428_c0_g1_NA.pdf) | -4.99 | 4.86e-04 | 5.11e-02 |
| isogroup51541_c0_g3K02A2_6-like | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup51541_c0_g3K02 | -4.68 | 4.23e-04 | 4.59e-02 |
| isogroup140490_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup140490_c0_g1_NA | -4.52 | 1.92e-05 | 9.28e-03 |
| isogroup65473_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup65473_c0_g1_NA.pdf) | -4.08 | 1.87e-04 | 2.42e-02 |
| isogroup3413_c0_g3_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3413_c0_g3_NA.pdf) | -3.71 | 7.52e-04 | 7.08e-02 |
| isogroup7218_c2_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7218_c2_g2_NA.pdf) | -3.01 | 7.26e-04 | 6.97e-02 |
| isogroup779_c0_g2CAAX-box_protein_maturation | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup779_c0_g2_CAAX | 2.90 | 1.92e-04 | 2.46e-02 |
| isogroup9138_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9138_c0_g2_NA.pdf) | 3.51 | 1.56e-04 | 2.19e-02 |
| isogroup1501_c0_g1von_Willebrand_factorvWFty | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup1501_c0_g1von | 4.56 | 1.65e-04 | 2.22e-02 |
| isogroup7806_c0_g1positive_regulation_of_CREB_tr | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7806_c0_g1_positi | 5.15 | 1.38e-04 | 2.18e-02 |
| isogroup14766_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14766_c0_g1_NA.pdf) | 5.53 | 7.43e-04 | 7.05e-02 |
| isogroup8406_c0_g1K02A2_6-like | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8406_c0_g1K02A | 6.28 | 7.36e-04 | 7.04e-02 |
| isogroup3220_c0_g3_DCUN1D3_cullin_family_protein_b | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3220_c0_g3_DCUN | 6.31 | 3.28e-04 | 3.76e-02 |
| isogroup13206_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13206_c0_g1_NA.pdf) | 6.58 | | 8.37e-02 |
| isogroup30129_c0_g5_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup30129_c0_g5_NA.pdf) | 6.62 | | 9.69e-02 |
| isogroup733_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup733_c0_g1_NA.pdf) | 6.73 | | 1.32e-03 |
| isogroup16276_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16276_c0_g1_NA.pdf) | 6.85 | | 4.04e-02 |
| isogroup34208_c0_g1_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup34208_c0_g1_NA.pdf) | 6.99 | | 2.22e-02 |
| isogroup5099_c0_g2_NA | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5099_c0_g2_NA.pdf) | 6.99 | | 8.37e-02 |
| isogroup3009_c0_g1_IRAK4_interleukin-1-mediated_si | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3009_c0_g1_IRAK4 | 7.01 | 1.16e-05 | 6.59e-03 |

From

| | From | From | From | | |
|--|------------|----------|-------|---|----------|
| Obe Import Import <th>to</th> <th>to</th> <th>to</th> <th></th> <th></th> | to | to | to | | |
| | Adjusted | | logFC | Image | ID |
| | 5.34e-02 | | 7 52 | ogreunt/1/268 c2 a1 N∆ | |
| Inspects Sporge_Invoked_analysis_with_DESeq2_And_all_databooplot.isogroup3332_c1_g2_post. 61.6 5.45-eV | 4 6.96e-02 | 7.22e-04 | | o- | isogroup |
| Integroup1733_c0_g3_NA | 4 6.62e-02 | 6.73e-04 | 8.08 | ogroup13196_c0_g3_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup13196_c0_g3_NA.pdf) | isogroup |
| InguresHS_sponge_RNAseq_analysis_with_DESeq2_AM_all_databoxplot.laogroup18022_c0_g1_NA | 5.52e-02 | 5.43e-04 | 8.16 | ogroup30332_c1_g2_positive_regulation_of_CREB_t (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup30332_c1_g2_posi | isogroup |
| (figuresHS_sponge_RNAseq_analysis_with_DESeq2_ANt_all_databoxplot.isogroup18022_c0_g1_NA pdf) 8.24 6.86 eAt | 5 8.78e-03 | 1.74e-05 | 8.19 | (liguresh5_sporige_knAseq_analysis_witit_bleseq2_Avin_aii_data/boxplot.isogroup755_c0_g5_nA.pdr) | isogroup |
| isogroup6115_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup6116_c0_g1_NA.pdf) 8.33 4.01e-04 isogroup67800_c1_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup67000_c1_g1_NA.pdf) 8.39 7.70e-04 isogroup6303_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup67000_c1_g1_NA.pdf) 9.23 5.02e-04 isogroup6305_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup6330_c0_g1_NA.pdf) 9.40 6.76e-04 isogroup6256_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup621_c0_g1_NA.pdf) 10.20 6.76e-04 isogroup12556_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup621_c0_g1_NA.pdf) 10.20 6.76e-04 isogroup12556_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup61_c0_g1_NA.pdf) 10.20 6.76e-04 isogroup12556_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup61_c0_g1_NA.pdf) 10.20 6.76e-04 isogroup12556_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup10756_c0_g1_NA.pdf) 10.20 6.76e-04 isogroup12556_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup10756_c0_g1_NA.pdf) 10.20 6.76e-04 isogroup1303_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup10756_c0_g1_NA.pdf) 10.20 6.77e-06 isogroup1303_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup10756_c0_g1_NA.pdf) 10.20 6.77e-06 isogroup1003_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup10756_c0_g1_NA.pdf) 10.20 6.77e-06 isogroup1003_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup1003_c0_g1_NA.pdf) 10.20 6.77e-06 isogroup1003_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogroup1003_c0_g1_NA.pdf) 10.20 6.77e-06 isogroup1003_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AW1_all_databoxplot.isogr | 4 6.15e-02 | 6.17e-04 | 8.20 | ogroup1148_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup1148_c0_g1_NA.pdf) | isogroup |
| isogroup67080_e1_g1_NA iggroup630_e0_g2_CNTN4_Contactin_4 iggroup630_e0_g2_CNTN4_Contactin_4 iggroup638_e0_g1_NA iggroup638_e0_g1_N | 4 6.59e-02 | 6.68e-04 | 8.24 | ogroup18022_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup18022_c0_g1_NA.pdf) | isogroup |
| isogroup2830_c0_g2_CNTN4_Contactin_4 isogroup8830_c0_g1_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup8836_c0_g1_NA.pdf) isogroup821_c0_g3_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup8636_c0_g1_NA.pdf) isogroup91_c0_g1_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup621_c0_g3_NA.pdf isogroup12556_c0_g1_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup621_c0_g3_NA.pdf isogroup12556_c0_g1_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup12556_c0_g1_NA.pdf) isogroup16037_c0_g1_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup12556_c0_g1_NA.pdf) isogroup16037_c0_g1_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup16037_c0_g1_NA.pdf) isogroup16037_c0_g1_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup16037_c0_g1_NA.pdf) isogroup19324_c0_g4_NA ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AVN_all_data/boxplot.isogroup848_c0_g1_Jerky | 4.41e-02 | 4.01e-04 | 8.33 | ogroup5115_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5115_c0_g1_NA.pdf) | isogroup |
| isogroup8636_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8636_c0_g1_NA.pdf) 9.54 6.76e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8621_c0_g3_NA.pdf) 9.54 6.76e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup81_c0_g1_NA.pdf) 10.00 4.07e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup91_c0_g1_NA.pdf) 10.00 4.07e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup10037_c0_g1_NA.pdf) 10.00 4.07e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup100324_c0_g4_NA.pdf) 10.00 4.07e-04 4.00 4.00 4.00 4.00 4.00 4.00 4.00 4 | 7.19e-02 | 7.70e-04 | 8.39 | (liguresho_sporige_kivAseq_arialysis_witri_bbseqz_Aviv_aii_data/boxplot.isogroups/060_c1_g1_ivA.pdi) | isogroup |
| isogroup621_c0_g3_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup621_c0_g3_NA.pdf) isogroup91_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup621_c0_g3_NA.pdf) isogroup12556_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup91_c0_g1_NA.pdf) isogroup16037_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12556_c0_g1_NA.pdf) isogroup16037_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16037_c0_g1_NA.pdf) isogroup14075_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14075_c0_g1_NA.pdf) isogroup848_c0_g1Jerky_protein_homolog-like (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup848_c0_g1Jerky 10.80 8.77e-08 isogroup19093_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_c0_g1_NA.pdf) isogroup19093_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_c0_g1_NA.pdf) isogroup90926_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_c0_g1_NA.pdf) isogroup90926_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_c0_g1_NA.pdf) isogroup90926_c0_g1_regulation_of_milotic_cell_cyc (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup90926_c0_g1_nA.pdf) isogroup90926_c0_g1_regulation_of_milotic_cell_cyc (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup90926_c0_g1_regul 12.00 8.29e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9090926_c0_g1_nA.pdf) 12.00 8.29e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9090926_c0_g1_nA.pdf) 12.00 8.29e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9090926_c0_g1_nA.pdf) 12.00 8.29e-04 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/ | 3 9.83e-02 | 1.15e-03 | 8.98 | (liguresH5_sponge_RNAseq_analysis_with_DESeq2_AVN_aii_data/boxplot.isogroup2830_c0_g2_CNTN | isogroup |
| isogroup91_c0_g1_NA isogroup12556_c0_g1_NA isogroup12556_c0_g1_NA isogroup16037_c0_g1_NA isogroup16037_c0_g1_NA isogroup16037_c0_g1_NA isogroup14075_c0_g1_NA isogroup14075_c0_g1_NA isogroup14075_c0_g1_NA isogroup14075_c0_g1_NA isogroup19324_c0_g1_NA isogroup1 | 5.23e-02 | 5.02e-04 | 9.23 | ogroup8636_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup8636_c0_g1_NA.pdf) | isogroup |
| isogroup12556_cO_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12556_cO_g1_NA.pdf) isogroup16037_cO_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16037_cO_g1_NA.pdf) isogroup14075_cO_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16037_cO_g1_NA.pdf) isogroup848_cO_g1_Jerky_protein_homolog-like (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup848_cO_g1_Jerky 10.80 8.77e-06 isogroup19032_d_cO_g4_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup848_cO_g1_Jerky 11.80 3.29e-05 isogroup19093_cO_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_cO_g1_NA.pdf) isogroup7076_cO_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_cO_g1_NA.pdf) isogroup7076_cO_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7076_cO_g1_NA.pdf) isogroup9226_cO_g1_regulation_of_mitotic_cell_cyc (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_cO_g1_regul 12.30 2.13e-05 isogroup9154_cO_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3154_cO_g1_MAP1 13.20 2.26e-04 | 4 6.62e-02 | 6.76e-04 | 9.54 | ogroup621_c0_g3_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup621_c0_g3_NA.pdf) | isogroup |
| isogroup16037_c0_g1_NA isogroup14075_c0_g1_NA isogroup14075_c0_g1_NA isogroup14075_c0_g1_NA isogroup14075_c0_g1_NA isogroup19324_c0_g4_NA isogroup19324_c0_g4_NA isogroup19324_c0_g4_NA isogroup19324_c0_g4_NA isogroup19093_c0_g1_NA | 4.46e-02 | 4.07e-04 | 10.10 | ogroup91_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup91_c0_g1_NA.pdf) | isogroup |
| isogroup14075_c0_g1_NA isogroup848_c0_g1_Jerky_protein_homolog-like isogroup19324_c0_g4_NA isogroup1993_c0_g1_NA isogroup19093_c0_g1_NA isogroup7076_c0_g1_NA isogroup9226_c0_g1_regulation_of_mitotic_cell_cyc isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9154_c0_g1_MAP1A. isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9154_c0_g1_MAP1A. isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9154_c0_g1_MAP1A. isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9154_c0_g1_MAP1A. isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr | 1 5.52e-02 | 5.44e-04 | 10.20 | ogroup12556_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup12556_c0_g1_NA.pdf) | isogroup |
| isogroup19324_c0_g4_NA isogroup1903_c0_g1_NA isogroup7076_c0_g1_NA isogroup7076_c0_g1_NA isogroup9326_c0_g1_regulation_of_mitotic_cell_cyc isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9154_c0_g1_MAP1 10.80 isogroup19324_c0_g4_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19324_c0_g4_NA.pdf) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_c0_g1_NA.pdf) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul 12.30 isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul 13.20 2.26e-04 | 4 6.02e-02 | 6.02e-04 | 10.50 | ogroup16037_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup16037_c0_g1_NA.pdf) | isogroup |
| isogroup19324_c0_g4_NA isogroup1903_c0_g1_NA isogroup7076_c0_g1_NA isogroup7076_c0_g1_NA isogroup9226_c0_g1_regulation_of_mitotic_cell_cyc isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9254_c0_g1_MAP1 ifiguresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup19093_c0_g1_NA.pdf) isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr | 5.08e-02 | 4.81e-04 | 10.80 | ogroup14075_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup14075_c0_g1_NA.pdf) | isogroup |
| isogroup19093_c0_g1_NA isogroup7076_c0_g1_NA isogroup9226_c0_g1_regulation_of_mitotic_cell_cyc isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup92154_c0_g1_MAP1) 13.20 2.26e-04 | 5.59e-03 | 8.77e-06 | 10.80 | ogroup848_c0_g1Jerky_protein_homolog-like (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup848_c0_g1Jerky | isogroup |
| isogroup19093_c0_g1_NA isogroup7076_c0_g1_NA isogroup9226_c0_g1_regulation_of_mitotic_cell_cyc isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup925_c0_g1_nAp1) i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup925_c0_g1_regul i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup925_c0_g1_regul i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup925_c0_g1_regul i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1 i(figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup93154_c0_g1_MAP1) | 5 1.26e-02 | 3.29e-05 | 11.60 | (TiguresHS_sponge_RNAseq_analysis_witn_DESeq2_AvN_ail_data/boxplot.isogroup19324_c0_g4_NA.pdf) | isogroup |
| isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul) 12.30 2.13e-05 (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup923154_c0_g1_MAP1) 13.20 2.26e-04 | 7.59e-02 | 8.29e-04 | 12.00 | pareup19093 c0 g1 NA | isogroup |
| isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup92154_c0_g1_MAP1 13.20 2.26e-04 | 7 9.05e-04 | 8.23e-07 | 12.20 | ogroup7076_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup7076_c0_g1_NA.pdf) | isogroup |
| isogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3154_c0_g1_MAP1 13.20 2.26e-04 | 5 9.74e-03 | 2.13e-05 | 12.30 | (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup9226_c0_g1_regul | isogroup |
| 0 | 1 2.78e-02 | 2.26e-04 | 13.20 | ogroup3154_c0_g1_MAP1A_Microtubule-associated_pr (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup3154_c0_g1_MAP1 | isogroup |
| | 7.19e-02 | 7.73e-04 | 15.00 | percupi102645 c0 a1 NA | isogroup |
| isogroup20684_c0_g3_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup20684_c0_g3_NA.pdf) 18.10 1.54e-06 | 3 1.49e-03 | 1.54e-06 | 18.10 | ogroup20684_c0_g3_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup20684_c0_g3_NA.pdf) | isogroup |
| isogroup23305_c0_g4_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup23305_c0_g4_NA.pdf) 20.60 3.56e-04 | 4.03e-02 | 3.56e-04 | 20.60 | ogroup23305_c0_g4_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup23305_c0_g4_NA.pdf) | isogroup |
| isogroup5533_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5533_c0_g1_NA.pdf) 24.20 2.71e-05 | 5 1.09e-02 | 2.71e-05 | 24.20 | ogroup5533_c0_g1_NA (figuresHS_sponge_RNAseq_analysis_with_DESeq2_AvN_all_data/boxplot.isogroup5533_c0_g1_NA.pdf) | isogroup |

← Previous Next →

(Page generated on Fri May 27 11:26:27 2022 by ReportingTools (http://research-pub.gene.com/ReportingTools/) 2.32.1 and hwriter (https://cran.r-project.org/web/packages/hwriter/index.html))